

IFW16

PATENT APPLICATION: US/09/930,559B

DATE: 10/01/2004 TIME: 10:33:30

Input Set : D:\ARCD351.APP.txt

Output Set: N:\CRF4\10012004\I930559B.raw

3 <110> APPLICANT: DAWSON, GLYN SEUNGUEN, JULIA CHO 6 <120> TITLE OF INVENTION: COMPOUNDS THAT ENHANCE TUMOR DEATH 8 <130> FILE REFERENCE: ARCD:351US 10 <140> CURRENT APPLICATION NUMBER: 09/930,559B 11 <141> CURRENT FILING DATE: 2001-08-15 13 <150> PRIOR APPLICATION NUMBER: 60/225,526 14 <151> PRIOR FILING DATE: 2000-08-15 16 <160> NUMBER OF SEQ ID NOS: 13 18 <170> SOFTWARE: PatentIn Ver. 2.1 20 <210> SEO ID NO: 1 21 <211> LENGTH: 2279 22 <212> TYPE: DNA 23 <213 > ORGANISM: Human 25 <400> SEQUENCE: 1 26 ggcacgagcg aagatggcgt cgcccggctg cctgtggctc ttggctgtgg ctctcctgcc 60 27 atggaeetge getteteggg egetgeagea tetggaeeeg eeggegeege tgeegttggt 120 28 gatctggcat gggatgggag acagctgttg caatccctta agcatgggtg ctattaaaaa 180 29 aatggtggag aagaaaatac ctggaattta cgtcttatct ttagagattg ggaagaccct 240 30 gatggaggac gtggagaaca gcttcttctt gaatgtcaat tcccaagtaa caacagtgtg 300 31 teaggeactt getaaggate etaaattgea geaaggetae aatgetatgg gatteteeca 360 32 qqqaqqccaa tttctqaqqq caqtqqctca gagatqccct tcacctccca tgatcaatct 420 33 gatetegqtt qqqqqacaac ateaaggtgt ttttggaete eetegatgee caggagagag 480 34 ctctcacatc tgtgacttca tccgaaaaac actgaatgct ggggcgtact ccaaagttgt 540 35 teaggaaege etegtgeaag eegaataetg geatgaeeee ataaaggagg atgtgtateg 600 36 caaccacagc atcttcttgg cagatataaa tcaggagcgg ggtatcaatg agtcctacaa 660 37 gaaaaacctg atggccctga agaagtttgt gatggtgaaa ttcctcaatg attccattgt 720 38 ggaccetgta gatteggagt ggtttggatt ttacagaagt ggecaageea aggaaaceat 780 39 tecettacag gagacetece tgtacacaca ggacegeetg gggetaaagg aaatggacaa 840 40 tgcaggacag ctagtgtttc tggctacaga aggggaccat cttcagttgt ctgaagaatg 900 41 gttttatgcc cacatcatac cattccttgg atgaaacccg tatagttcac aatagagctc 960 42 agggagccc taactettee aaaccacatg ggagacagtt teetteatge eeaageetga 1020 43 getcagatec agettgeaac taateettet ateatetaac atgeactact tggaaagate 1080 44 taagatetga atettateet ttgccatett etgttaceat atggtgttga atgcaagttt 1140 45 aattaccatg gagattgttt tacaaacttt tgatgtggtc aagttcagtt ttagaaaagg 1200 46 gagtetgtte cagateaggg ceagaaetgt geeeaggeee aaaggagaea aetaaetaaa 1260 47 gtagtgagat agattetaag ggcaaacatt tttccaagte ttgccatatt tcaagcaaag 1320 48 aggtgcccag gcctgaggta ctcacataaa tgctttgttt tgctggtgat ttaaccagtg 1380 49 cttggaaaaa tcttgcttgg ctatttctgc atcatttctt aaggctgcct tcctcttga 1440 50 gtacgttgcc ctctgtgcta tcaatcatct tatcatcaat tattagacaa atcccactgg 1500 51 cctacagtet tgettetgea geacecaett tgteteetea ggtagtgatg aattagttge 1560 52 tgtcacaaaa ggagggaagt agcacccaaa ttaaattgct taagagagga aatgtacatc 1620 53 ttgtataact tagggagcga agaaaatgta ggcgcgaaag tgaaaagtga ggcagctagt 1680

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55 tagagtcaac ttactctgtt gctggcttta gcagagaata ggaggaacca tatgaaaaag 1800
56 atcaggettt etgaetteca tececaaaac acatttaeca geataeteca aactgtttet 1860
57 gatgtgttcc atgagaaaag gattgtttgc tcaaaaagct tggaaaatac tacacactcc 1920
58 ctttctcctt ctggagatca acccacatta gagtgtctaa ggactcctga gaattcctqt 1980
59 tacagtaaac aaaactaacg taatctacca tttcctacac tatttgagca tggaaatcat 2040
60 agtccccact ctatgaaaac ttaacgcttt ttggaagaca tttctgtagc atgtcagttt 2100
61 ggagaaatga tgagctacgc cttgatgaaa gaaccgtgtt ggtgctgcta agtttagcca 2160
62 ttatggtttt tcctttctct ctcttaagcc ttattcttca actaaaagat gaggattaag 2220
63 agcaagaagt tgggggggat gtgaaaataa ttttatgagg ttgtctaaaa tctcgtgcc 2279
66 <210> SEQ ID NO: 2
67 <211> LENGTH: 306
68 <212> TYPE: PRT
69 <213> ORGANISM: Human
71 <400> SEQUENCE: 2
72 Met Ala Ser Pro Gly Cys Leu Trp Leu Leu Ala Val Ala Leu Leu Pro
                    5
                                        10
75 Trp Thr Cys Ala Ser Arg Ala Leu Gln His Leu Asp Pro Pro Ala Pro
76
                20 '
78 Leu Pro Leu Val Ile Trp His Gly Met Gly Asp Ser Cys Cys Asn Pro
79
            35
                                40
81 Leu Ser Met Gly Ala Ile Lys Lys Met Val Glu Lys Lys Ile Pro Gly
84 Ile Tyr Val Leu Ser Leu Glu Ile Gly Lys Thr Leu Met Glu Asp Val
                        70
87 Glu Asn Ser Phe Phe Leu Asn Val Asn Ser Gln Val Thr Thr Val Cys
                    85
90 Gln Ala Leu Ala Lys Asp Pro Lys Leu Gln Gln Gly Tyr Asn Ala Met
               100
                                   105
93 Gly Phe Ser Gln Gly Gly Gln Phe Leu Arg Ala Val Ala Gln Arg Cys
                               120
96 Pro Ser Pro Pro Met Ile Asn Leu Ile Ser Val Gly Gly Gln His Gln
                           135
99 Gly Val Phe Gly Leu Pro Arg Cys Pro Gly Glu Ser Ser His Ile Cys
                        150
102 Asp Phe Ile Arg Lys Thr Leu Asn Ala Gly Ala Tyr Ser Lys Val Val
                    165
                                        170
105 Gln Glu Arg Leu Val Gln Ala Glu Tyr Trp His Asp Pro Ile Lys Glu
106
                180
                                    185
108 Asp Val Tyr Arg Asn His Ser Ile Phe Leu Ala Asp Ile Asn Gln Glu
                                200
                                                    205
111 Arg Gly Ile Asn Glu Ser Tyr Lys Lys Asn Leu Met Ala Leu Lys Lys
                            215
114 Phe Val Met Val Lys Phe Leu Asn Asp Ser Ile Val Asp Pro Val Asp
115 225
                        230
                                            235
117 Ser Glu Trp Phe Gly Phe Tyr Arg Ser Gly Gln Ala Lys Glu Thr Ile
                    245
                                        250
120 Pro Leu Gln Glu Thr Ser Leu Tyr Thr Gln Asp Arg Leu Gly Leu Lys
                                    265
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123 Glu Met Asp Asn Ala Gly Gln Leu Val Phe Leu Ala Thr Glu Gly Asp
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126 His Leu Gln Leu Ser Glu Glu Trp Phe Tyr Ala His Ile Ile Pro Phe
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                            295
129 Leu Gly
130 305
133 <210> SEQ ID NO: 3
134 <211> LENGTH: 7
135 <212> TYPE: PRT
136 <213> ORGANISM: Human
138 <400> SEQUENCE: 3
139 Gly Cys Val Lys Ile Lys Lys
140 1
143 <210> SEQ ID NO: 4
144 <211> LENGTH: 8
145 <212> TYPE: PRT
146 <213> ORGANISM: Human
148 <400> SEQUENCE: 4
149 Ile Arg Tyr Cys Trp Leu Arg Arg
150
    1
                     5
153 <210> SEQ ID NO: 5
154 <211> LENGTH: 9
155 <212> TYPE: PRT
156 <213> ORGANISM: Human
158 <400> SEQUENCE: 5
159 Val Thr Thr Leu Cys Cys Gly Lys Asn
160 1
163 <210> SEQ ID NO: 6
164 <211> LENGTH: 7
165 <212> TYPE: PRT
166 <213> ORGANISM: Human
168 <400> SEQUENCE: 6
169 Met Leu Cys Cys Met Arg Arg
170 1
173 <210> SEQ ID NO: 7
174 <211> LENGTH: 8
175 <212> TYPE: PRT
176 <213> ORGANISM: Human
178 <400> SEQUENCE: 7
179 Met Gly Cys Leu Gly Asn Ser Lys
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183 <210> SEQ ID NO: 8
184 <211> LENGTH: 8
185 <212> TYPE: PRT
186 <213> ORGANISM: Human
188 <400> SEQUENCE: 8
189 Met Gly Cys Leu Gly Asn Ser Lys
190 1
193 <210> SEQ ID NO: 9
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194 <211> LENGTH: 10
195 <212> TYPE: PRT
196 <213> ORGANISM: Human
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200
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203 <210> SEQ ID NO: 10
204 <211> LENGTH: 35
205 <212> TYPE: DNA
206 <213> ORGANISM: Artificial Sequence
208 <220> FEATURE:
209 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
210
          Primer
212 <400> SEQUENCE: 10
213 totaggtacc aagatggcgt cgcccggctg cctgt
                                                                       35
216 <210> SEQ ID NO: 11
217 <211> LENGTH: 38
218 <212> TYPE: DNA
219 <213> ORGANISM: Artificial Sequence
221 <220> FEATURE:
222 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
223
          Primer
225 <400> SEQUENCE: 11
226 acggtctaga tcatccaagg aatggtatga tgtqqqca
                                                                       38
229 <210> SEQ ID NO: 12
230 <211> LENGTH: 5
231 <212> TYPE: PRT
232 <213> ORGANISM: Artificial Sequence
234 <220> FEATURE:
235 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
236 Peptide
238 <400> SEQUENCE: 12
239 Val Lys Ile Lys Lys
240 1
243 <210> SEQ ID NO: 13
244 <211> LENGTH: 5
245 <212> TYPE: PRT
246 <213> ORGANISM: Artificial Sequence
248 <220> FEATURE:
249 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
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252 <400> SEQUENCE: 13
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254 1

VERIFICATION SUMMARY

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